

**Amendments to the Specification:**

Please replace the paragraph beginning at page 6, line 15, with the following rewritten paragraph:

-- Particularly conserved regions and amino acid residues common to nectin polypeptides were identified by aligning nectin polypeptide sequences with each other and additional closely-related members of the nectin-Ig superfamily of proteins. The amino acid sequence of nectin-3 $\alpha$  and nectin-4 (SEQ ID Nos: 6 and 24) were compared with the amino acid sequences of other nectin and Ig family members (SEQ ID NO:20, 22, and 25), using a multiple sequence alignment program. The alignment of these sequences is shown in Table 2, and includes consensus residues (capitalized), which are identical among at least a majority (three) of the five amino acid sequences in the alignment. In addition, lower case residues are shown on a separate line of Table 2 and represent residues that are not consensus residues, but are identical between human nectin-3 $\alpha$  and human nectin-4 (SEQ ID Nos: 6 and 24). --

Please replace Table 2 beginning at page 7, line 7, with the following rewritten Table:

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Table 2  
Conserved Nectin Amino Acids

(Hs=Homo sapiens)

(Mus=Murine)

Protein (SEQ ID NO)					
HUNECTIN2 ([SEQ ID NO:]22)	~~~~~	~~~~~	MARAAALLPS	RSPPTPLLWP	LLLLLLL...
HUCD155 ([SEQ ID NO:]25)	~~~~~	~~~~~	~~~~~	MARAMAAAWP	LLLVALLVLS
HUNECTIN1 ([SEQ ID NO:]20)	~~~~~	~~~~~	~~~~~MARMG	LAGAAGRWWG	L...ALGLTA
HUNECTIN3 ([SEQ ID NO:]6)	MARTLRPSPL	CPGGGKAQLS	SASLLGAGLL	LQPPTPPPLL	LLLFPLLFS
HUNECTIN4 ([SEQ ID NO:]24)	~~~~~	~~~~~	~~~~~MPLSLG	AEMWGPEAWL	LLLLLLASFT
consensus				<u>P</u> <u>W</u>	LLL LL <u>f</u>
				<u>l</u>	<u>l</u>
HUNECTIN2	51				100
HUCD155	..ETGAQDVR	VQVLPEVRGQ	LGGTVELPCH	L.LPPVPGLY	ISLVTWQRPD
HUNECTIN1	WPPPGTGDVV	VQAPTQVPGF	LGDSVTLPCY	LQVPNMEVTH	VSQLTWAR..
HUNECTIN3	FFLPGVHSQV	VQVNDSMYGF	IGTDVVLHCS	FANP.LPSVK	ITQVTWQK.S
HUNECTIN4	RLCGALAGP.	IIVEPHVTAV	WGKNVSLKCL	I..EV..NET	ITQISWEKIH
	GRCP..AGE.	LETSDVVTVV	LGQDAKLPCF	YRGDS..GEQ	VGQVAWARVD
	[ cPG ag	VQV VtGv	LG V LPC	P e	I QV W R]
	<u>PG</u>	<u>VQV</u> <u>V G</u>	<u>LG</u> <u>V LPC</u>	<u>P</u>	<u>I QVTW R</u>
	<u>c</u> <u>ag</u>	<u>t v</u>		<u>e</u>	
HUNECTIN2	101				150
HUCD155	APANHQNVAA	FHPKMGPSFP	SPKPGSERLS	FVSAKQSTGQ	DTEAELQDAT
HUNECTIN1	.HGESGSMVA	FHOTQGPSYS	E....SKRLE	FVAARLG...	...AELRNAS
HUNECTIN3	TNGSKQNVAI	YNPSMGVSV.	.LAPYRERVE	FL.....	..RPSFTDGT
HUNECTIN4	.GKSSQTVAV	HHPQYGFSVQ	..GEYQGRVL	FKNYSLN...	.....DAT
	AGEGAQELAL	LHSKYGLHVS	..PAYEGRVE	QPPPPRNPL.	.....DGS
	[g Q A	H yG SV	Y gRVE	F n	DAT]
	Q VA	HP G SV	Y RVE	F	DAT
	g	y	g	n	

HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	151 LALHGLTVED EGNYTCEFAT FPKGSVRGMT WLRVIAKPKN QAEAQKVTF. LRMFGLRVED EGNYTCLFVT FPQGSRSVDI WLRVLAKPQN TAEVQKVQL. IRLSRLELED EGVYICEFAT FPTGNRESQL NLTVMKPTN WIEGTQAVLR ITLHNIGFSD SGKYICKAVT FPLGNAQSST TVTVLVEPTV SLIKGPDSLI VLLRNAVQAD EGEYECRVST FPAGSFQARL RLRVLVPPLP SLNPGP.ALE [ L nL ED EG Y C F T FP GS q LRVLAKP N s E L] L L ED EG Y C F T FP GS LRVLAKP N E L n q v sl gp	200
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	201 ....SQDPTT VALCISKEGR PPARISWLSS LDWEAKETQV SGTLAGTVTV ....TGEPVP MARCVSTGGR PPAQITWHSD LGGMPNTSQV PGFLSGTVTV AKKGQDDKVL VATCTSANGK PPSVVSWE TR LKGEARVPGD SGTPMAPVTV DGGNE...TV AAICIAATGK PVAHIDWEGD LGEM..ESTT TSFPNETATI EGQGL...TL AASC.TAEGS PAPSVTWDTE VKGT..TSSR SFKHSRSAAV [g T aA C Sa G PPA I W L G S SG TVTV] T A C SA G PPA I W L G S SG TVTV g a a	250
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	251 TSRFTLVPSG RADGVTVTCK VEH..ESFEE PALIPVTL SV RYPPEVSISG TSLWILVPSS QVDGKNVTCK VEH..ESFEK PQLLTVNLT V YYPEVSISG ISRYRLVPSR EAHQOSLACI VNYHMDRFKE ....SLTLNV QYEPEVTIEG ISQYKLFPTR FARGRRITCV VKHP..ALEK DIRYSFILDI QYAPEVSVTG TSEFHLVPSR SMNGQPLTCV VSHP..GLLQ DQRITHILHV SFLAEASVRG [TS LVPSR A G TC V Hp FE d r iL V Y PEVSI G] TS LVPSR A G TC V H FE L V Y PEVSI G v p l d r i v	300
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	301 Y.DDN.WYLG RTDATLSCDV RSNPEPTGYD WSTTSGTFPT SAVAQGSQVL Y.DNN.WYLG QNEATLTCDA RSNPEPTGYN WSTTMGPLPP FAVAQGAQLL F.DGN.WYLG RMDVKLTCKA DANPPATEYH WTTLNGSLPK GVEAQNRTLF Y.DGN.WFVG RKGVNLCNA DANPPPFKSV WSRLDGQWPD GLLASDNTLH LEDQNLWHIG REGAMLCCLS EGQPPPSYN. WTRLDGPLPS GVRVDGDTLG [YD N WYLG R gA LkC A NPPPTY WSTLdG LP G AQG TL] Y D N WYLG R A L C A NPPPT Y WSTL G LP G AOG TL g k r d	350
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	351 IH.AVDSL FN TTFVCTVTNA VGMGRAEQVI FVRETP.... IR.PVDKPIN TTLICNV TNA LGARQAE LTV QVKEGP.... FKGPINYS LA GTYICEATNP IGTRSGQVEV NITEFPYTPS FVHPLTFNYS GVIYICKVTNS LGQRSDQKVI YISDPPTTTT LQPTIQWHPS F.PPLTTEHS GIYVCHVSNE FSSRDSQVTV DVLDPQEDSG KQ..... [F Plt s G YIC VTN G R Q V EpP q] F P GTYIC VTN G R Q V V E P lt s dp g	400
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	401 .....RAS P...RDV..G PLVWGAVGGT LLVLLLLLAGG .....PSE H...SGISRN AIIFLVLG.. ILVFLILLGI .....PPE HGRRAGPVPT AIIGGVAGSI LLVLIIVGGI TADIEDLATE PKKLPFPLST LATIKDDTIA TIIASVVGGA LFIVLVSVLA ....VDLV... SAS VVVVGVI AAL LFCLLVVVVV [ d II GV G LLVLLV vG] II GV G LLVLLV G dl f v	450
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	451 SLAFILLRVR RR.....RKS .PGGAGGGAS GDGGFYDPKA QVLGNGDPVF GIYFYWSKCS REVLWHCHLC .PSSEHHQSC RN~~~~~ VVALRRRRHT FKGDYSTKKH .VYNGYSKA GIPQHHPMA QNLQYPDDSD GIFCYRRRT FRGDYFAKNY IPPSDMQKES QIDVLQODEL D..SYP.DSV LMSRYHRR...KAQQMTQKY EEELTLTREN SIRRLHSHHT DPRSQPEESV [ y RR y P e I lH d Ls PD Sv] Y RR P I P S y e l d s v	500

HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	501 WTPVVPGPME P.DGKDEEEE EEEKAEEKGL MLPPPPALED DMESQLDGSL ~~~~~ .DEKKAGPLG G.SSYEEEEEE EEEGGGGGER KVGGPHPKYD EDAKRPYFTV .KKENKNP.. .VNNLIRKDY LEEPEKTQWN NVENLNRFER PMDYEDLKM GLRAEGHPDS LKDNSSCSVM SEEPEGRSYS TLTTVREIET QT...ELLSP [ p n Eepe e 1] P EE E n pe e 1	550
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	551 ISRRVYV~ ~~~~~ ~~~~~ DEAEARQDGY GDRTLGYQYD PEQLDLAENM VSQNDGSFIS KKEWYV~ GM.KFVSDEH YDENEDDLVS HV...DGSVI SR...REWYV ~~~~~ GSGRAEEEEED QDEGIKQAMN HFVQENGTLR AKPTGNGIYI NGRGHLV [g A E DE I H g y] A D g e e h g y	597